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Perfect score:
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match Length
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| (cgn2_6/ptodata/1/ina/6A_COMB.seq:*
| (cgn2_6/ptodata/1/ina/6A_COMB.seq:*
| (cgn2_6/ptodata/1/ina/6B_COMB.seq:*
| (cgn2_6/ptodata/1/ina/backfīles1.seq:*
| (cgn2_6/ptodata/1/ina/backfīles1.seq:*
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Copyright (c) 1993 - 2005
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US-09-433-699-13-1560
US-09-270-767-15231
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US-09-902-540-1360
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US-09-902-540-13617
US-09-903-16-12617
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US-09-671-325-1570
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31566, Ap
31560, Ap
31501, Ap
1196, Ap
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1570, Ap
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Patent No. 6639063
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Property Application NUMBER: US/09/621,976
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
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Sequence 2892,	Patent No.	Patent No.	Sequence	Sequence	Sequence	Sequence	Seguence	Sequence	Seguence	Sequence	Sequence	Sequence	Seguence	Sequence	Sequence	Sequence	Sequence
2892, Ap	5386025	5386025	Appli	4, Appli	2, Appli	2, Appli	Appli	5, Appli	1143, Ap	11, Appl	15, Appl	986, App	8129, Ap	4163, Ap	8972, Ap	4648, Ap	4444, Ap

ALIGNMENTS

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RESULT 2
US-09-513-999C-1128/c
US-09-513-999C-1128/c
Sequence 1128, Application US/09513999C
Patent No. 6783961
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclett, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 13204
LENGTH: 442
TYPE: DNA
ORGANISM: Homo sapiens
US-09-621-976-13204
                                                         SOFTWARE: Patent.pm
SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100
Matches 20; Conservative
                                                                                            NUMBER OF SEQ ID NOS: 36681
ORGANISM: Homo sapiens
                                     ENGTH: 467
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; LOCATION: 162..
US-09-513-999C-1128
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                                                                         NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4169

LENGTH: 2527

TYPE: DNA

ORGANISM: Human

S-09-949-016-4169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/09433699B
Patent No. 6165786
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF NUCLEOLIN EXPRESSION
FILE REFERENCE: RTG-0109
CURRENT APPLICATION NUMBER: US/09/433,699B
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 3
LENGTH: 2518
TYPE: NNS
                                         Query Match
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Best Local (
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEC TO MOC.
                                                                                                                                                                                                                                                                                                                                              APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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-09-433-699-3
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NAME/KEY: CDS
LOCATION: (112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
   Local Similarity 100 nes 20; Conservative
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o. 6812339
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Pred. No. 0.055;
Mismatches 0
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OF DETECTION
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-433-699-14
RESULT 7
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; ORGANISM: Human
US-09-949-016-15911
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US-09-949-016-15911/c
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CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: ANTISENSE MODULATION OF NUCLEOLIN EXPRESSION FILE REFERENCE: RTS-01.09
CURRENT APPLICATION NUMBER: US/09/433,699B
CURRENT APPLICATION 1999-11-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 14
                                                                                                                                 Matches
                                                                                                                                                 Query Match
Best Local :
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Patent No. 6165786
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Best Local
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J.
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: C. Frank Bennett APPLICANT: Lex M. Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                  LENGTH: 20
TYPE: DNA
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                                                                                                                                                 Similarity
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Pred. No.
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Pred. No.
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0.78;
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US-09-134-000C-2181

Sequence 2181, Application US/09134000C Patent No. 6617156

GENERAL INFORMATION:

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US-09-270-767-31566
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PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 2620
LENGTH: 1176
Type: No.
                                                                                                      Patent No. 6703491

GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
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                                                                    SOFTWARE: PatentIn Ver. SEQ ID NO 31566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local (
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Best Local
                                                                                                                                                                                                                                                                 Sequence 31566, Application US/09270767
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -09-489-039A-2620
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Klebsiella pneumoniae
                  ORGANISM: Drosophila melanogaster
                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Enterococcus faecalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                      ENGTH: 1211
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Query Match

75.0%; Score 15;

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Length 1211;

Hinkle, Slater,

, Gregory J. , Steven C. d, Roger C.

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RESULT 12
US-09-902-540-3960
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US-09-489-039A-2450/c
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                 Sequence 3960, Application US/09902540 Patent No. 6833447 GENERAL INFORMATION:
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Patent No. 6610836
GENERAL INFORMATION:
APPLICANT: Gary Breton et. al
                                                                                                                                                                                                                                                                                                                          SEQ ID NO 15231
LENGTH: 1498
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LENGTH: 1395
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Best Local
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                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
APPLICANT: Goldman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 14342
                                                                                                                                                                                                                                                                                         ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                                                                                                                           TYPE: DNA
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   Barry S.
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APPLICANT: Goldman, Barry S.

APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

APPLICANT: Wiegand, Roger C.

ITILE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15849)8

CURRENT APPLICATION NUMBER: US/09/902,540

CURRENT FILING DATE: 2001-07-10

PRIOR APPLICATION NUMBER: 60/217,883

PRIOR FILING DATE: 2000-07-10

NUMBER OF SEQ ID NOS: 16825

SEQ ID NO 1196

LENGTH: 18324

TYPE: DNA

ORGANISM: Myxococcus xanthus

US-09-902-540-1196
                   GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

PRIOR PRICATION NUMBER: 60/237,768

PRIOR PILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR SEQ. ID NOS: 207012
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US-09-949-016-12617/c
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US-09-902-540-3960
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Best Local Similarity luv.
Thes 15; Conservative
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LENGTH: 2118
TYPE: DNA
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Best Local (
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
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CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
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OF DETECTION
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Search completed: April 12, 2005, 01:20:40 Job time : 128 secs

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PRIOR FILING DATE: 2000-10-20
PRIOR PELICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-09
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, NAME/KEY: misc feature
; LOCATION: (1)...(160018)
; OTHER INFORMATION: n = A,T,C
US-09-949-016-12617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-949-016-15994/c
                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 15994
                                                                 Matches
                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15994, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES,
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
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Best Local :
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                            FEATURE: misc_feature NAME/KEY: misc_feature LOCATION: (1)...(160018) OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                LENGTH: 160018
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/ 75.0%; Score 15;
Local Similarity 100.0%; Pred. No.
15; Conservative 0; Mismatcl
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OF DETECTION AND USES THEREOF
                                                                                           Length 160018;
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Minimum DB
Maximum DB
                                                                                                                                                                                                            Result
No.
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Word size
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OM nucleic - nucleic search, using sw model
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                                                     0000000
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                                                                                                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      seq
     NNNNNNNNNNN
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length: 2000000000
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20
                                                                                                                                                                                                            Query
Match
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   1000.0
1000.0
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238.583 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5615251 seqs, 3030001701 residues
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(cgn2 6/ptodata/2/pubpna/US07 PUBCOMB.seq:*

(cgn2 - 6/ptodata/2/pubpna/US07 PUBCOMB.seq:*
                                                                                                                                                                                                                                                                                                                                                          /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US108_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US108_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US106_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version Copyright (c) 1993 - 2005
                                                                                                                                                                                                              Length
   240
377
396
438
444
636
1276
1276
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US-10-797-822-1
US-10-377-930-35092
US-10-357-930-35092
US-09-918-995-37497
US-10-357-930-4798
US-10-357-930-13957
US-10-198-846-9321
US-10-363-345A-27853
US-10-363-345A-27853
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Compugen Ltd.
                                                                                                                                                                                                            Description
Sequence 1, Appli
Sequence 2, Appli
Sequence 35092, A
Sequence 37497, A
Sequence 4788, Ap
Sequence 43939, A
Sequence 13957, A
Sequence 27853, A
Sequence 27853, A
Sequence 27853, A
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3	3	3	1276	-	115-10-363-4834-27854
13	200	100.0	1860	18	-10-357-9
14	20	8	2518	9	09-825-886
15	20	8	2518	18	83-
16	20	•	2637	17	-10-104-047-
17	20	•	2723	17	-10-384-569-5
18	20	•	3201	18	-10-357-930
19	20	•	3201	18	-10-357-
20	20	•	3497	14	-10-198-846
21	20	•	6112	15	-10-172-086
22	20	•	6112	18	-10-311-507-
23	20	•	6112	18	-10-480-846-3
24	20	•	10942	9	09-825
25	20	100.0	10942	17	US-10-384-569-4
26	18	•	1212	18	-10-43
27	16	•	378	18	-10-437-963-12
28	16	•	422	17	-10-424-599-
29	15	٠	198	18	-10-437-963-9
30	15	٠	279	18	-10-437-963-392
31	15	•	468	18	-10-43
32	15	•	516	15	-10-156-761-238
33	15	75.0	622	17	-10-425-114
34	15	•	657	18	-10-437-
35	15	•	688	17	-10-425-114
36	15	•	693	17	-10-425-114
37	15	•	693	18	-425-115
38	15	•	702	18	-10-425-115-113
39	15	•	725	18	-10-363-3452
40	15	75.0	725	18	-10-363-345A-14
41	15	•	725	19	-10-363-483A-14
42	15	•	725	19	-10-363-483A-14
43	15	•	726	18	-10-363-345A-20
44	15	•	726	18	-10-363-345A-20
45	15	٠			

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ALIGNMENTS

Sequence 1, Application US/10797822
Publication No. US20050026860A1
GENERAL INFORMATION:
APPLICANT: Chin-Tarng Lin
APPLICANT: Chin-Tarng Lin
APPLICANT: Han-Chung Wu
TITLE OF INVENTION: Nucleolin Antisense Sequence FOR Inhibition of CANCER Cell Prolif
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/10/797,822
CURRENT FILING DATE: 2004-03-09
NUMBER OF SEQ ID NOS: 2
SEQ ID NO 1
SEQ ID NO 1

RESULT 1 US-10-797-822-1

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; LENGTH: 20
; TYPE: DNA
; ORGANISM: Artificial Sequence
; ORGANISM: INFORMATION: Antisense fr.
; OTHER INFORMATION: nucleolin mR.
US-10-797-822-1
RESULT 2
US-10-797-822-2/c
; Sequence 2, Application US/10797822
; Publication No. US20050026860A1
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                                                                                                                                                 Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                          ATGATGGCGGCGGAGTGTGA 20
                                                                                           ATGATGGCGGCGGAGTGTGA 20
                                                                                                                                                     100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                            Antisense fragment which sequence is specifically complementary nucleolin \ensuremath{\mathsf{mRNA}}
                                                                                                                                                     0
                                                                                                                                                     Score 20; DB
Pred. No. 0.1
0; Mismatches
                                                                                                                                                                                 DB 19;
                                                                                                                                                        0
                                                                                                                                                                                 Length
                                                                                                                                                        Indels
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RESULT 4
US-09-918-995-37497/C
, Sequence 37497, Application US/09918995
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                                                                                                                                                                                                                                                                                            SEQ ID NO 35092
LENGTH: 240
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 2
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 62232
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
APPLICANT: Monahan, John
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 2

EQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/189,862
PRIOR FILING DATE: 2000-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/357,930 CURRENT FILING DATE: 2003-02-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/10/797,822
CURRENT FILING DATE: 2004-03-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Chin-Tarng Lin
APPLICANT: Han-Chung Wu
TITLE OF INVENTION: Nucleolin Antisense Sequence FOR Inhibition of CANCER Cell Prolif
                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/785,276
PRIOR FILING DATE: 2003-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LO-357-930-35092/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Sense fragment which sequence is similar to the nucleolin mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 2000-05-25
APPLICATION NUMBER: 60/211,314
FILING DATE: 2000-06-09
APPLICATION NUMBER: 60/219,007
FILING DATE: 2000-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/207,454
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                                                                                                                                                         1 ATGATGGCGGCGGAGTGTGA 20
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                                                                                                                                                                                                                    Similarity
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                                                                                                                    ATGATGGCGGCGGAGTGTGA 97
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                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                  Score 20; DB 1
Pred. No. 0.08;
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                                                                                                                                                                                                                                    DB 18;
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GENERAL INFORMATION:

APPLICANT: Hyseq, Inc.

TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED

TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES

FILE REFERENCE: 20411-756

CURRENT APPLICATION NUMBER: US/09/918,995

CURRENT FILING DATE: 2001-07-30

PRIOR APPLICATION NUMBER: US/09/235,076

PRIOR FILING DATE: 1999-01-20

NUMBER OF SEQ ID NOS: 38054

SOFTWARE: FRASISEQ for Windows Version 3.0

SEQ ID NO 37497

LENCTH: 377
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                                                                                                                                            US-10-357-930-4788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA; ORGANISM: Homo sapiens US-09-918-995-37497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: MRI-007BCN
CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: 09/785,276
PRIOR FILING DATE: 2003-02-16
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/189,862
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR APPLICATION NUMBER: 60/207,454
                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4788
                                                             Matches
                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MODAHAN, JOHN
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS,
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENTITLE OF INVENTION: HUMAN PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Schlegel. Robert APPLICANT: Endege, Wilson
                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 62232
                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2000-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/219,007
                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-06-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/211,314
                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: 224, 242, 273, 286, 312,
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2000-05-25
                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98 ATGATGGCGGCGGAGTGTGA 79
                                                                                                                                                                                                                                                                               396
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                                                                              Similarity
ATGATGGCGGCGGAGTGTGA 20
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                                                                              100.0%;
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Pred. No. 0.
                                                                              Score 20; DB 18
Pred. No. 0.074;
                                                             Mismatches
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                                                                                                DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 10;
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T, PREVENTION, AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                  Length 396;
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                                                             Gaps
                                                             0
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ATGATGGCGGCGGAGTGTGA 67

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Sequence 13957, Application US/10357930

Publication No. US20040259086A1

GENERAL INFORMATION:

APPLICANT: Schlegel. Robert

APPLICANT: Schlegel. Robert

APPLICANT: Monahan, John

ITILE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR

TITLE OF INVENTION: HUMAN PROSTATE CANCER

FILE REFERENCE: MRI-007BCN

CURRENT APPLICATION NUMBER: US/10/357,930

CURRENT APPLICATION NUMBER: 09/785,276

PRIOR APPLICATION NUMBER: 00/183,319

PRIOR APPLICATION NUMBER: 60/183,319

PRIOR APPLICATION NUMBER: 60/183,319

PRIOR FILING DATE: 2000-02-16

PRIOR APPLICATION NUMBER: 60/183,319

PRIOR APPLICATION NUMBER: 60/207,454

PRIOR APPLICATION NUMBER: 60/207,454

PRIOR FILING DATE: 2000-05-25

PRIOR APPLICATION NUMBER: 60/211,314
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/785,276
PRIOR FILING DATE: 2003-02-16
PRIOR PELICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/189,862
PRIOR FILING DATE: 2000-03-16
PRIOR PELING DATE: 2000-03-16
PRIOR PELING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR PILING DATE: 2000-05-25
PRIOR PILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR PILING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR PILING DATE: 2000-07-18
PRIOR PILING DATE: 2000-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 62232
SOFTWARE: PRECSEQ for Windows Version 4.0
SEQ ID NO 43939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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APPLICANT: Endege, Wilson
APPLICANT: Endege, Wilson
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR
TITLE OF INVENTION: LIGHTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF
TITLE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
CURRENT FILING DATE: 2003-02-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 438
TYPE: DNA
APPLICATION NUMBER: 60/211,314 FILING DATE: 2000-06-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/255,281 FILING DATE: 2000-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    80 ATGATGGCGGCGGAGTGTGA 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20;
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100.0%; Pred. No. 0.073;
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APPLICANT: STEALMANN, KATCHIEGO
APPLICANT: STEALMANN, KATCHIEGO
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND
TITLE OF INVENTION: FOR IDENTIFICATION, ASSESSMENT, PRE
TITLE OF INVENTION: THERAPY OF BREAST CANCER
FILE REFERENCE: MRI-049
CURRENT APPLICATION NUMBER: US/10/198,846
CURRENT FILING DATE: 2002-07-18
CURRENT FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/306,220
PRIOR FILING DATE: 2001-07-18
NUMBER OF SEQ ID NOS: 14084
SOSTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 9321
LENGTH: 636
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, NAME/KEY: misc feature
; LOCATION: 3, 9, 56, 59
; OTHER INFORMATION: n = A,T,C or
US-10-198-846-9321
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US-10-198-846-9321/c
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                                                                                                                                                                                                                                                                                                                     RESULT 9
Sequence 27853, Application US/10363345A
Publication No. US20040234960A1
GENERAL INFORMATION:
APPLICANT: Alexander Olek
APPLICANT: Kurt Berlin
TITLE OF INVENTION: Method for determining the degree of methylation of defined
TITLE OF INVENTION: Cytosines in genomic DNA in the sequence context of 5'-CpG-3
FILE REFERENCE: E011227
CURRENT APPLICATION NUMBER: US/10/363,345A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
SEQ ID NO 27853
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; SEQ ID NO 13957
; LENGTH: 444
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Best Local Similarity 100.0%;
Matches 20; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9321, Application US/10198846 Publication No. US20030099974A1 GENERAL INFORMATION:
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Best Local Similarity 100.0%; Pred. No. 0.072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lillie, James APPLICANT: Xu, Yongyao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 62232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
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Wang, Youzhen
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Pred. No. 0.068;
0; Mismatches
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PREVENTION,
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US-10-363-483A-27853
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US-10-363-345A-27854/c
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US-10-363-345A-27853
                                                                                                                                                                                                                                                                                                                                                  US-10-363-483A-27853
                                                                                                                                                                                                                                                                                             Sequence 27853, Application US/10363483A Publication No. US20050064401A1 GENERAL INFORMATION:
                                                                                                 SEQ ID NO 27853
LENGTH: 1276
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LENGTH: 1276
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Best Local :
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                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/363,483A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
                                                                                                                                                                                    APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
APPLICANT: Kurt Berlin
TITLE OF INVENTION: Diagnosis of illnesses or
TITLE OF INVENTION: illnesses
FILE REFERENCE: 82011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
APPLICANT: Christian Piepenbrock
APPLICANT: Kurt Berlin
TITLE OF INVENTION: Method for determining the degree of methylation of defined
TITLE OF INVENTION: cytosines in genomic DNA in the sequence context of 5'-CpG-
FILE REFERENCE: E01/1227
CURRENT FILING DATE: 2003-03-03
UNMBER OF SEQ ID NOS: 40712
              OTHER INFORMATION: OTHER INFORMATION:
                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                                                                                  TYPE: DNA
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            chemically
CpG-island
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            treated genomic DNA (Homo sapiens) No: 27853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treated genomic DNA No: 27854
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 20; DB 18;
Pred. No. 0.061;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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Query Match

100.0%;

Score 20;

DB 19;

Length 1276;

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US-10-363-483A-27854/c
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LENGTH: 1276
TYPE: DNA
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 62232
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 28900
LENGTH: 1860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 28900, Application US/10357930 Publication No. US20040259086A1
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Best Local (
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Matches 20; Conservative (
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CURRENT APPLICATION NUMBER: US/10/363,483A
CURRENT FILING DATE: 2003-03-03
NUMBER OF SEQ ID NOS: 40712
                                                                                                                                       PRIOR APPLICATION NUMBER: 60/211,314
PRIOR TILING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR FILING DATE: 2000-07-18
                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/189,862
PRIOR FILING DATE: 2000-03-16
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILING DATE: 2000-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: MODAMAN, JOHN
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Diagnosis of illnesses or predisposition to certain TITLE OF INVENTION: illnesses
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APPLICANT: Christian Piep
APPLICANT: Kurt Berlin
                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: 09/785,276
PRIOR FILING DATE: 2003-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
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les 20; Conserv
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APPLICANT: RUGSlahti, Erkki
APPLICANT: Porkka, Kimmo
APPLICANT: Porkka, Kimmo
APPLICANT: Christian, Sven
ITILE OF INVENTION: HMGNZ Peptides and Related Molecules
ITITLE OF INVENTION: that Selectively Home to Tumor Blood Ve
FILE REFERENCE: P-LJ 5662
CURRENT APPLICATION NUMBER: US/10/400,083
CURRENT FILING DATE: 2003-03-20
PRIOR APPLICATION NUMBER: US 10/116,866
PRIOR APPLICATION NUMBER: US 0/16,866
PRIOR FILING DATE: 2002-04-05
NUMBER OF SEO ID NOS: 20
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 18
IENGTH. 2518
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APPLICANT: Muller, Sylviane
APPLICANT: Muller, Sylviane
APPLICANT: Muller, Sylviane
APPLICANT: Muller, Sean-Paul
APPLICANT: Griend, Giles
TITLE OF INVENTION: A NOVEL CELL SURFACE RECEPTOR FOR HIV RETROVIRUSES,
TITLE OF INVENTION: THERAPBUTIC AND DIAGNOSTIC USES.
FILE REFERENCE: 03495.0166-01.000
CURRENT APPLICATION NUMBER: US/09/825,886
CURRENT FILING DATE: 2001-07-26
PRIOR APPLICATION NUMBER: 09/393,302
PRIOR APPLICATION NUMBER: 09/393,302
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: 09/393,002
PRIOR FILING DATE: 1999-03-12
PRIOR FILING DATE: 1999-03-12
PRIOR APPLICATION NUMBER: 06/040,969
PRIOR APPLICATION NUMBER: 60/040,969
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; ORGANISM: Homo sapiens
US-09-825-886-28
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US-10-357-930-28900
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NUMBER OF SEQ ID NOS: 32
SOFTWARE: PatentIn Ver. 2
SEQ ID NO 28
LENGTH: 2518
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Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 20; Conservative 0; Mismatches
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Publication No. US20040186056A1
                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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APPLICANT: Callebaut,
TYPE: DNA
ORGANISM: Homo sapiens
                                                        ENGTH: 2518
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Krust, Bernard
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: yongsung@mail.kribb.re.kr
Plate: 9 row: G column: 05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genome Research Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kim, Y.S.
21C Frontier Korean EST Project 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.R., Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                       Similarity
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    ATGATGGCGGCGGAGTGTGA 111
                              ATGATGGCGGCGGAGTGTGA 20
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ilarity 100.0%;
Conservative C
                                                                               100.0%;
larity 100.0%;
Conservative (
                                                                                                                                                                            /note=Worgan: Cervix; Vector: pCNS-D2; Site 1: ECORI; Site 2: NotL; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoRI site by treatment of T4 RNA ligase and the first strand CDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. Coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of the cDNA libraries constructed by this method are full-length enriched cDNA library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="Uterine"
/cell_type="Epithelial"
/cell_line="SNU-17"
/lab_host="Top10F'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="C1SNU17-9-G05"
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Pred. No. 0.34;
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Pred. No. 0.34;
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RESULT 4
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AUTHORS
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ORGANISM
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VERSION
KEYWORDS
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AL121075/c
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Best Local Similarity
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          TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                      59
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 160)

Dias Neto, E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H. Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. an Simpson,A.J.
                                                                                                                                                                                                                                                                                       160 bp mRNA linear
QVO-CT0583-101100-491-a04 CT0583 Homo sapiens cDNA,
BQ319320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No s1 sequence available.
This clone (DKrZp762B035) is available at the RZPD in Berlin.
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany This is the 5' sequence of the clone insert Clone from S. Wiemann, Molecular Genome Analysis, German Cancer Research Center (DKF2); Email s.wiemann@dkfz- heidelberg.de; sequenced by GBF (National Research Centre for Biotechnology Ltd., Braunschweig/Germany) within the cDNA sequencing consortium of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria; Prim.
1 (bases 1 to 155)
Bloecker, H., Boecher, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         155 bp mRNA linear EST 04-S
DKFZp762B035 r1 762 (synonym: hmel2) Homo sapiens cDNA clone
DKFZp762B035-5', mRNA sequence.
                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                     BQ319320.1 GI:20925236 EST.
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Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="DKFZp762B035"
/clone="DKFZp762B035"
/tissue_type="melanoma (MeWo cell line/dev stage="adult"
/lab host="DH10B"
/clone_lib="762 (synonym: hmel2)"
/note="Vector: pSport1; Site_1: Not1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 0.34;
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                                                    de Souza, S.J. and
                                                                                                                                                                                                                                                                                                                mRNA sequence.
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                                                                       Simpson, D.H.,
                                                                                                                              Briones, M.R.,
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sequence tags

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REFERENCE
AUTHORS
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Best Local
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                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 163)

Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Ragai, M.A., da Silva, W. Jr., Zago, M.A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., D'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                          Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  163 bp mRNA linear
RC2-HT0861-310700-011-f06 HT0861 Homo sapiens cDNA,
BE719607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fax: +55-11-270/vv.
Email: asimpson@ludwig.org.br
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human
This sequence was derived from the following URL
This sequence was derived from the following URL
This sequence was derived from the FAPESP/LICR Human
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                                                                                                                                                                                                     Shotgun sequencing of the human transcriptome with ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=QV0&t2=QV0-CT0583-101100-491-a04&t3=2000-11-10&t4=1)
Seq_primer: puc 18 forward
                                                                                                                                                                                                                           Simpson, A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                    sequence tags
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Ludwig Institute for Cancer Research
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                                                                                                                         10737800
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                                    Prof. Antonio
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           quality sequence start: 55 quality sequence stop: 95. Location/Qualifiers
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+55-11-2704922
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/mol_type="mRNA"
/db_xref="taxon:9606"
/dev_stage="Adult"
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                                    Prudente 109,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Li,Y., Xu,C., Fang,R., Guegler,K., Rao,M.S., Murage,J., Fisk,G.J., Li,Y., Xu,C., Fang,R., Guegler,K., Rao,M.S., Mandalam,R., Lebkowski,J and Stanton,L.W.

Transcriptome characterization elucidates signaling networks that control human ES cell growth and differentiation
Nat. Biotechnol. 22 (6), 707-716 (2004)

Contact: Brandenberger R
                                                                                                                                                                                                                                                                                                                                                          230 Constitution Drive, Menlo Park, Tel: 650 473 8658 Fax: 650 473 7760
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1 (bases 1 to 164)
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Email: asimpson@ludwig.org.br
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=RC2-HT0861-310
700-011-f066t3=2000-07-31&t4=1)
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Location/Qualifiers
                                                                                                                                                                                                                                                                                    Email: rbrandenberger@geron.com
Insert Length: 164 Std Error:
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Regenerative Medicine
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                                                               /clone_lib="GRN_ES"
/note="oligo dT primed, full-length enriched cDNA library
/note="oligo dT primed, full-length enriched cDNA library
from undifferentiated hES cell lines H1 (p22), H7 (p29),
and H9 (p26) maintained in feeder-free conditions"
                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/dev_stage="Adult"
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RESULT 8
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                                                                                                                                                                                              20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=MRO-NN0087-230
500-009-d06&t3=2000-05-33&t4=1)
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Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V.,
O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
Simpson,A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence tags
Proc. Natl. Acad. Sci. U.S.A.
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MRO-NN0087-230500-009-d06
BE699907
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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llarity 100.0%;
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/clone_lib="NN0087"
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/mol_type="mRNA"
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                                                                                                        AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: cgapbs r@mail.nih.gov
Oligo-dT track not found, Not I site shown in beginning of
is likely internal to the message. Tissue Procurement: Loui
Staudt, M.D., Ph.D.
CDNA Library Preparation: M.B. Soares Lab
CDNA Library Arrayed by: M.B. Soares Lab
CDNA Library Arrayed by: M.B. Soares Lab
CDNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
www.bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 77
Statistical analysis of the 5' untranslated region of human mRNA using 'Oligo-Capped' cDNA libraries Genomics 64 (3), 286-297 (2000)
                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Hom 1 (bases 1 to 185)
Suzuki, Y., Ishihara, D., Sasaki, M., Nakagawa, H., Hata, H., Tsunoda, T., Watanabe, M., Komatsu, T., Ota, T., Isogai, T., and Sugano, S.
                                                                                                                                                                                                                                                mRNA sequence.
AU077224
AU077224.1 GI:7439847
                                                                                                                                                                                                                                                                                                           AU077224 Sugano cDNA library Homo sapiens cDNA clone 2 similar to 5'-end region of Chinese hamster nucleolin
                                                                                                                                                                                      Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
1 (bases 1 to 184)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMAGE:3062234 3', mRNA sequence.
AW575670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UI-HF-BM0-adk-h-02-0-UI.sl NIH_MGC_38 Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: M13 Forward
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AW575670.1 GI:7247209
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /Clone="IMAGE:3062234"
/tissue_type="lymph"
/tissue_type="lymph"
/cell_type="germinal center B cells"
/cell_type="germinal center B cells"
/cell_line="MGC85"
/lab host="MHC85"
/clone_lib="NHH MGC 38"
/clone_lib="NHH MGC 38"
/note="Vector: p7773-Pac; Site_1: NotI; Site_2: Eco RI;
/note="Vector: p7773-Pac; Site_1: NotI; Site_2: Eco RI;
Constructed from size fractionated cytoplasmic mRNA
(2.5-3.5kb). Directionally cloned. Cells provided by Louis
M. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58
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Louis M.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              137
                                                                                                                                                                                                                    found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
rms-rt Length: 2161 Std Error: 0.00
Seq primer: -40Up from Gibco
High quality sequence stop: 171.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AW050536 205 bp mRNA linear EST 09-MAR-2000 wz14a01.x1 NCI CGAP Ut4 Homo sapiens cDNA clone IMAGE:2557992 3' similar to gb:M60858_rna1 NUCLEOLIN (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ATGATGGCGGCGGAGTGTGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki.Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and
Sugano,S. Construction and characterization of a full
length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone was obtained from a '5'-end-enriched' cDNA library constructed by 'Oligo-Capping' method. The coding region starts from the 50 bp upstream to the 3'-end.

Location/Qualifiers
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National Cancer Institute, Cancer Genome Anat
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumor Gene Index
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Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                      CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                    /clone="IMAGE:2557992"
/tissue_type="serous papillary
pooled tumors"
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/db_xref="taxon:9606"
/clone="Zrv61966"
/clone_lib="Sugano cDNA library"
                                                                                                                                                      /organism="Homo sapiens"
/mol_type="mRNA"
/lab_host="DH10B"
/clone_lib="NCI_CGAP_Ut4"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: SalI;
                                                                                                                                    /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Homo sapiens"
|mol_type="mRNA"
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Pred. No.
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                                                                                           carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ph.D.,
                                                                                         high grade,
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AUTHORS
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SOURCE
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W79364
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 210)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M
Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 ATGATGGCGGCGGAGTGTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: mob.REGA+ET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 722 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Washington University
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                              /dev_stage="19 weeks"
/lab_host="pH10B (ampicillin resistant)"
/lab_host="pH10B (ampicillin resistant)"
/clome_lib="Soares fetal_heart_NbHH19W"
/clome_lib="Soares fetal_heart_nbHH19W"
/clome_lib="Soares fetal_heart_nbH19W"
/clome_lib="Soares fetal_hear
                             normalization to a Cot = 5. Library constructed by M.Fatima Bonaldo. This library was constructed from the same fetus as the fetal lung library, Soares fetal lung NbHL19W."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.48 kb. Life Technologies catalog #:
11542-016"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="GDB:1271572"
/db_xref="taxon:9606"
/clone="IMAGE:346197"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Homo sapiens"
|mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sex="unknown"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   School of Medicine way, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 20; DB
Pred. No. 0.3
0; Mismatches
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Query Match Best Local Similarity

100.0%; ilarity 100.0%; Conservative

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Score 20; DB 7; Pred. No. 0.34; Mismatches C

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Indels

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Gaps

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Length 210;

Matches

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KEYWORDS
SOURCE
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BX644913/c
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AA263089/c
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                     ORGANISM
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                                                                                                                                                                                                                                                   85
                                                                                                         BX644913 219 bp DKFZp781I1236 r1 781 (synonym: h DKFZp781I1236 5', mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claudio, J.O., Liew, C.C., Dempsey, A.A., Cukerman, E., Stewart, J. Na, E., Atkins, H.I., Iscove, N.N. and Hawley, R.G.
Identification of sequence-tagged transcripts differentially expressed within the human hematopoietic hierarchy
Genomics 50 (1), 44-52 (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The Toronto Hospital
CRCS-424, 67 College St., Toronto,
Tel: 416 3403834
Fax: 416 3403453
                     Homo sapiens
                                                                          BX644913.1 GI:34479246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KG1a primary library.
Seq primer: 5' GAAATTAACCCTCACTAAAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: r.hawley@utoronto.ca
Similar to M60858 nucleolin gene. Clone was randomly picked from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oncology Research Laboratories
The Toronto Hospital
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Hawley RG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 215)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA sequence.
AA263089
AA263089.1 GI:1898887
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PMY0457 KG1-a Lambda Zap
                                   iomo sapiens (human)
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                                                                                                                                                                                                                                               ATGATGGCGGCGGAGTGTGA 66
                                                                                                                                                                                                                                                                                  ATGATGGCGGCGGAGTGTGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATGATGGCGGCGGAGTGTGA
                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                              /clone lib="KGi-a Lambda Zap Express cDNA library" /note="Vector: Lambda Zap Express (Stratagene); Site_1: EcoRI; Site_2: XhoI; Unidirectional cloning sites: EcoRI-XhoI _mRNA was purified from KGI-a cell line, cDNA was synthesized using an XhoI-OligodT linker primer. EcoRI adaptors were ligated, followed by digestion with XhoI for directional cloning into predigested Lambda Zap Express"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /cell_type="promyeloblast"
/cell_line="KG1-a"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
 Metazoa; Chordata; Craniata;
                                                                                                                                                                                                                                                                                                                                       100.0%; Score 20; DB 1; Length 215; 100.0%; Pred. No. 0.34;
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cDNA library Homo
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Vertebrata; Euteleostomi;
                                                                                                                               sapiens
                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                 CDNA clone
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RESULT 14
BF757253
LOCUS
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MEDLINE
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
Email: asimpson@ludwig.org.br
This sequence was derived from t
Project. This entry can be seen
                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 220)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Baia,G.S., Simpson,D.H., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., Brunstein,A., Geoliveira, P.S., Bucher, P., Jongeneel,C.V., Brunstein,A., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                          Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
                                                                Tel: +55-11-2704922
Fax: +55-11-2707001
                                                                                                            Brazi.
                                                                                                                                                                                          Contact: Simpson A.J.G.
                                                                                                                                                                                                                                                                                                   Shotgun sequencing of the human transcriptome with ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QV0-CT0583-031100-476-e03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  No s1 sequence available.
This clone (DKFZp78111236) is available at the RZPD in Berlin.
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 1405
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany This is the 5' sequence of the clone insert Clone from S. Wiemann, Molecular Genome Analysis, German Cancer Research Center (DKFZ); Email s.wiemann@dkfz- heidelberg.de; sequenced by AGGWA (Berlin/Germany) within the cDNA sequencing consortium of the German Genome Project.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 219)

Wambutt,R., Heubner,D., Mewes,H.W., Weil,B., Amid,C., Osanger,A., Fobo,G., Han,M. and Wiemann,S.

EST (Wambutt,R., Heubner,D., Mewes,H.W., Weil,B., Amid,C., et al. Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BF757253.1
                                                                                                                                                                                                                                                                               sequence tags
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/db xref="taxon:9606"
/clone="DKFZp78111236"
/dev stage="adult"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="781 (synonym: hlcc4)"
/note="Vector: pSport1_Sfi; Site_1:
cDNA-collection"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GI:12105153
                                                                                                                                                                                                                                                           Acad.
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  from the
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FAPESP/LICR Human the following URL
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                        Cancer Genome
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AI185752/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumor Gene Index
Unpublished (1997)
Unpublished (1997)
Contect: Robert Strausberg, Ph.D.
Contect: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
IMAGE Tanath: 954 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI185752 221 bp mRNA linear EST 29-OCT-1998 qe44e01.xl Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone: IMAGE:1741848 3' similar to gb:m60858_rna1 NUCLEOLIN (HUMAN);, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ب
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1 (Dases 1 to 221)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome.Anatomy Project (CGAP);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AI185752
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Seq primer: puc 18 forward
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Homo sapiens
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Location/Qualifiers
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    /dev_stage="19 weeks"
//lab_host="DH108 (ampicillin resistant)"
//clone_lib="Soares fetal lung_NbHL19W"
//clone_lib="Soares fetal lung_NbH19W"
//clone_lib="Soares fetal lung_NbH19W"
//clone_lib="Soares feta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=Torgan: colon; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1741848"
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Soares and M.Fatima Bonaldo. This library was constructed from the same fetus as the fetal heart library, Soares fetal heart NbHH19W."

 Matches Query Match Best Local Similarity 20; Conservative 100.0%; 0 Score 20; Pred. No. Mismatches 0.34; DB 1; Length 221 Indels <u>,</u>

Gaps

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ORIGIN

밁 ફ 37 1 ATGATGGCGGCGGAGTGTGA 20 ATGATGGCGGCGGAGTGTGA 18

Search completed: April 12, 2005, 01:18:23 Job time : 3107 secs

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Database :
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Perfect score:
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BD024875 Sequence
AJ33862 Homo sapi
AJ339439 Homo sapi
AJ342352 Homo sapi
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BC002343 Homo sapi
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ALIGNMENTS

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RESULT 2
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AUTHORS
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                                                                                                                                                                       116 ATGATGGCGGCGGAGTGTGA 97
                                            CQ472921 396 bp D
Sequence 4788 from Patent WOO160860.
CQ472921
CQ472921.1 GI:41438540
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Homo sapiens (human)
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Millennium Predictive Medicine, Inc. (U
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
Homo sapiens
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Sequence 35092 from Patent
CQ503225
CQ503225.1 GI:41468861
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Genes differentially expressed in human prostate cancer and their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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CQ512072/c
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AR421707/c
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Sequence 43939 from Patent WO
CQ512072.1 GI:41478336
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Millennium Predictive Medicine, Inc. (
Location/Qualifiers
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Mammalia; Eutheria;
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Sequence 13204 from patent U
AR421707
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Location/Qualifiers
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1 (bases 1 to 442)
Edwards,J.-B.D.M., Jobert,S. and Giordano,J.-Y.
EST's and encoded human proteins
Patent: US 6639063-A 13204 28-OCT-2003;
Location/Qualifiers
1. .442
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent: WO 0160860-A 43939 23-AUG-2001;
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larity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                      100.0%; Score 20; DB 6; llarity 100.0%; Pred. No. 3.7; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DN
/db_xref="taxon:9606"
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Primates;
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AX982401/c
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BD117260/c
LOCUS
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Matches 20
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Best Local Similarity
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                                                                                  Patent: JP 20
GENSET CORP
OS Homo sal
PN JP 2002
PD 15-JAN-
PF 07-AUG-
PR 05-AUG-
PI JEAN BA
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EST and encoded human protein.
BD117260
BD117260.1 GI:23212164
JP 2002010789-A/9337.
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1. (bases 1 to 442)
Edwards, J.B.D.M., Jobert, S. and Giordano, J.E.
EST and encoded human protein
Patent: JP 2002010789-A 9337 15-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dumas Milne Edwards, J.B., Jobert, S. and ESTs and encoded human proteins Patent: EP 1104808-A 13204 06-JUN-2001;
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C12N15/00
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C12N1/21
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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llarity 100.0%; Pred. No. 3.7
Conservative 0; Mismatches
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JP 2002010789-A/9337
15-JAN-2002
                        C12N5/10,C12P21/02,C12P21/08,C12Q1/68,C12N15/00,C12N5/00,
                                                      C12N15/09, C12N15/09, C07K14/47, C07K16/18, C12N1/15, C12N1/19,
                                                                                    07-AUG-2000 JP 2000280989
05-AUG-1999 US 60/147499
JEAN BAPUTIST DUWAS MILNE EDWARDS, SEVELIN JOBERT, JEAN EVE
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   and encoded
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/mol_type="unassigned DN
/db_xref="taxon:9606"
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CQ482090/c
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CQ482090
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                                                                     Dumas Milne Edwards, J.B., Duclert, A. and (
Expressed sequence tags and encoded human
Patent: EP 1033401-A 1128 06-SEP-2000;
                                                                                                                 Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                              Sequence 1128 from Patent
AX885265
AX885265.1 GI:40041253
                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent: WO 0160860-A 13957 23-AUG-2001;
Millennium Predictive Medicine, Inc. (U
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                        Homo sapiens (human)
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ilarity 100.0%; Pred. No. 3.7
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source
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/organism="Homo sapiens"
/mol_type="unassigned DNA"
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                                               ocation/Qualifiers
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                                                                                  and Giordano, J.Y.
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                                                                    RESULT 10
HSA338862/c
           ACCESSION
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           Homo sapiens
NL1-UP5C.
AJ338862
AJ338862.1 G
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PN JP 200
PD 02-001
PD 02-001
PF 24-FEE
PR 26-FEE
PR 26-FEE
PI JEAN (
PI JORDAN
PI JORDAN
PI C12N5/10,
PC C12P21
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 467)

Edwards, J. B.D.M., Duclair, B. and Jordan, J.Y.

Sequence tag and encoded human protein
Patent: JP 2001269182-A 1121 02-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence tag and encoded human BD024875
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                                                                                                   ATGATGGCGGCGGAGTGTGA 144
                                                                                                                                                                 100.0%;
ilarity 100.0%;
Conservative (
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26-FEB-1999 US 60/1224
JEAN BAPUTIST DUMAS MILNE
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JP 2001269182-A/1121
02-OCT-2001
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/proteIn_id="CAB99379.1"
/db_xref="G1:40041254"
/translation="MVKLAKAGKNQGDPKKMAPPPKEVEEDSEDEEMSEDEEDDSSGE
/translation="MVKLAKAGKNQGDPKKMAPPPKKVAAVTPGKKAAATPAKKTVTPAK
                                                                                                                                                                                                                                                                                                              Location/Qualifiers 162. .467.
Location/Qualifiers
                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/note="unnamed protein product"
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DUMAS MILNE EDWARDS,EIMERIC DUCLAIR,JEAN YVES
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sequence
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Pred. No. 3.7;
0; Mismatches
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Pred. No. 3.7
0; Mismatches
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HSA339439/c
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NR1-WA5C.
AJ339439
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(bases 1 to 620)

Kutsenko, A.S., Gizatullin, R.Z., Al-Amin, A.N., Wang, F., Kvasha, S.M., Podowaki, R.M., Matushkin, Y.G., Gyanchandani, A., Muravenko, O.V., Levitsky, V.G., Kolchanov, N.A., Protopopov, A.I., Kashuba, V.I., Kisselev, L.L., Wasserman, W., Wahlestedt, C. and Zabarovsky, E.R. NotI flanking sequences: a tool for gene discovery and verification of the human genome

Nucleic Acids Res. 30 (14), 3163-3170 (2002)
                                                                                                                                             Direct Submission
Direct Submission
Submitted (16-MAY-2001) Microbiology
Submitted (16-MAY-2001) Theorells vag,
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Direct Submission
Submitted (16-MAY-2001) Microbiology
Submitted Theritute, Theorells vag,
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Homo sapiens
Eukaryota; Metazoa;
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Kutsenko,A.S., Gizatullin,R.Z., Al-Amin,A.N., Wang,F., Kvasha,S.M., Podowski,R.M., Matushkin,Y.G., Gyanchandani,A., Muravenko,O.V., Levitsky,V.G., Kolchanov,N.A., Protopopov,A.I., Kashuba,V.I., Kisselev,L.L., Wasserman,W., Wahlestedt,C. and Zabarovsky,E.R. NotI flanking sequences: a tool for gene discovery and verification of the human genome
                                                                                                                                                                                                         Zabarovsky, E.R.
                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/clone="NL1-UP5C"
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                                    /mol_type="genomic DN/db_xref="taxon:9606"
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3, Box 280, Stockholm 171
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Box 280, Stockholm 171 77,
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RESULT 13
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1 (bases 1 to 716)

Rutsenko, A.S., Gizatullin, R.Z., Al-Amin, A.N., Wang, F., Kvasha, S.M., Podowski, R.M., Matushkin, Y.G., Gyanchandani, A., Muravenko, O.V., Levitsky, V.G., Kolchanov, N.A., Procopopov, A.I., Kashuba, V.I., Kisselev, L.L., Wasserman, W., Wahlestedt, C. and Zabarovsky, E.R., NotI flanking sequences: a tool for gene discovery and verification of the human genome
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.

1 (bases 1 to 788)

Kutsenko, A.S., Gizatullin, R.Z., Al-Amin, A.N., Wang, F., Kvasha, S.M., Podowski, R.M., Matushkin, Y.G., Gyanchandani, A., Muravenko, O.V., Levitsky, V.G., Kolchanov, N.A., Protopopov, A.I., Kashuba, V.I., Kisselev, L.L., Wasserman, W., Wahlestedt, C. and Zabarovsky, E.R., Notl flanking sequences: a tool for gene discovery and verification of the human genome
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                                                                                                                                       Homo sapiens
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Zabarovsky, E.R.
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Box 280, Stockholm 171
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1 (bases 1 to 863)

Kutsenko, A.S., Gizatullin, R.Z., Al-Amin, A.N., Wang, F., Kyasha, S.M., Podowski, R.M., Matushkin, Y.G., Gyanchandani, A., Muravenko, O.V., Levitsky, V.G., Kolchanov, N.A., Protopopov, A.I., Kashuba, V.I., Kisselev, L.L., Wasserman, W., Wahlestedt, C. and Zabarovsky, E.R. NotI flanking sequences: a tool for gene discovery and verification of the human genome

Nucleic Acids Res. 30 (14), 3163-3170 (2002)
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Direct Submission
Submitted (16-MAY-2001) Microbiology and Tumorbiology Centre,
Submitted (16-MAY-2001) Microbiology and Tumorbiology Centre,
Sarolinska Institute, Theorells vag, 3, Box 280, Stockholm 171 77,
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Location/Qualifiers
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Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Search completed: April 12, Job time : 1678 secs

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Abv43920 Human pro
Abv13966 Human pro
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Acn88171 Breast ca
Acg41262 Oligonucl
Abg41263 Oligonucl
Adg44512 Human col
Abv28882 Human pro
Adp44511 Human pro
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Aav71745 Human V3
Aac92553 Human nuc
Acc81089 Human nuc
Add49219 Human nuc
Adp54085 Human pRO
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ALIGNMENTS

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RESULT 1
ABV35074/c
ID ABV350
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XX ABV350
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XX Human;
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XX Homo s
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16-MAR-2000; 2000US-0189862P.
25-MAY-2000; 2000US-0207454P.
09-JUN-2000; 2000US-0211314P.
18-JUL-2000; 2000US-0219007P.
13-DEC-2000; 2000US-0255281P.
                                                                                                                           WPI; 2001-662795/76.
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                                                                                                                                                                                                                                                                                                                                                                                        cancer; marker;
                                                                                                                                                                           PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; carcinogen;
gene; ss.
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The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether

Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer.

Claim 1; Page 7324; 11750pp; English.

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RESULT 2
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                          The invention relates to an isolated polynucleotide comprising any one 38043 cDNA sequences, appearing as ACH12789-ACH50831, whose sequence wa determined by the technique of SBH (sequencing by hybridisation). Also included is a purified polypeptide comprising a sequence corresponding a reading frame of the novel polynucleotide. The nucleic acid sequences are useful in diagnostics as expressed sequence tags (EST) for identifying expressed genes or for physical mapping of the human genome in forensics, in assessing biodiversities, or in identifying mutations responsible for genetic disorders and other traits. The nucleotide sequences are also useful as hybridisation probes, as oligomers for CR for chromosome and gene mapping, in the recombinant production of the morarstina artisance number and professional professional properties of the markish professional professi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polynucleotide sequences obtained from various as hybridization probes, as oligomers for PCR, for mapping, in the recombinant production of protein,
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DICKSON M C.
JONES L W.
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is one of the 38043 isolated cDNA/EST sequences. No
for this patent did not form part of the printed to
obtained in electronic format directly from USPTO
seqdata.uspto.gov/sequence.html?DocID=20030073623
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16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
                                                                   The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the efficacy progression of prostate cancer in a patient; (c) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (f) assessing the prostate cancer has metastasized in a patient; (h) determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a patient.
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       396
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2000US-0207454P.
2000US-0211314P.
2000US-0219007P.
2000US-0255281P.
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       BP;
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                                                                                                                                                                                                                                                                                                                                                                824-825; 11750pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer; cytostatic; carcinogen; pharmacodyanamic
marker; gene; ss.
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         125
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                                                      85
    A; 94 C;
                                                      a pharmacodyanamic
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Pred. No.
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         <u>ი</u>
         83
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         T; 0 U;
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                                                      pharmacogenomic
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            6 Other;
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Query Match

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Best Local S
Matches 20
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16-MAR-2000; 2000US-0189862P.
25-MAY-2000; 2000US-0207454P.
09-JUN-2000; 2000US-0211314P.
18-JUL-2000; 2000US-0219007P.
13-DEC-2000; 2000US-0255281P.
                                                                                                                                                                   The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cell carcer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a patient; (II) is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                              Sequence 438
                                                                                                                                                                                                                                                                                                                                                                            Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef for detecting presence of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 8734; 11750pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schlegel R,
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                                                                                                        Similarity
                                                               ATGATGGCGGCGGAGTGTGA
                                                                                           Conservative
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marker; gene; ss.
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                                                                                                       100.0%;
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                                                                                                                                              A; 113 C; 118
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ABV13966/c
ID ABV13966 standard; cl
XX
AC ABV13966;
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DT 13-SEP-2002 (first |
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DE Human prostate expre
XX
Human prostate canc
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DA Human; prostate canc
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DA Human; prostate canc
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DA Homo sapiens.
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DA Homo sapiens.
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23-AUG-2001; 2001WO-
XX
DA 17-PEB-2000; 2000US-
PR 16-MAR-2000; 2000US-
PR 15-JUN-2000; 2000US-
PR 19-JUN-2000; 2000US-
PR 13-DEC-2000; 2000US-
PR (MILL-) MILLENNIUM F
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RESULT 6
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Best Local Similarity
Matches 20; Conser
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                        Human secreted protein 5'
                                                                                        06-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef for detecting presence of prostate cancer.
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16-MAR-2000; 2000US-0189862P.
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                                                                                                                                                                                                         AAC01130
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                                                                                                                                                                                                                                                                                                                                                                                           ATGATGGCGGCGGAGTGTGA
                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                         ATGATGGCGGCGGAGTGTGA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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; 2000US-0211314P.
; 2000US-0219007P.
; 2000US-0255281P.
                                                                                        (first entry)
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                              EST,
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Pred. No. 0.19;
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                                 SEQ
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RESULT 7
ACN88171/c
ID ACN8811
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XX ACN881
XX O2-DEC
XX Breast
XX Cancer
XX Cancer
XX US2003
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                   Cancer; breast; tumour; cytostatic; marker;
                                                                                                                                                                                  Breast cancer related marker,
                                                                                                                                                                                                              02-DEC-2004
                                                                                                                                                                                                                                                                        ACN88171 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 1128; 71pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dumas Milne Edwards
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      18-JUL-2001; 2001US-0306220P
                                18-JUL-2002; 2002US-00198846
                                                              29-MAY-2003
                                                                                          US2003099974-A1
                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               n; 5' EST;
therapy; c
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DB; AAG01124.
                                                                                                                                                                                                                                                                                                                                                 163
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                                                                                                                                                                                                                                                                                                                                                 ATGATGGCGGCGGAGTGTGA 144
                                                                                                                                                                                                                                                                                                                                                                  ATGATGGCGGCGGAGTGTGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  467 BP; 133 A; 127 C; 123 G; 84 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chromosome
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                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                              entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mapping;
                                                                                                                                                                                                                                                                           636 BP
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                                                                                                                                                                                     9321
                                                                                                                                                        detection; therapy;
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XX Uhuma
XX Grug
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated polypeptide (I) associated with a breast cancer which is encoded by a nucleic acid molecule comprising a nucleid sequence (S1). Further disclosed is an antibody that binds to the polypeptide of the invention. The activity of the polypeptide of the invention may be described as cytostatic. The antibody is useful for detecting the presence of (I) in a sample. Nucleic acid molecules of the invention are useful in the detection of breast tumours. (I) is useful as a marker for breast cancer and in breast cancer therapy. Sequences given in records ACN78851-ACN92934 represent nucleic acid markers associated with breast cancer. Note: The sequence listing does not form part of the specification but may be obtained in electronic format from the USPTO web site at seqdata.uspto.gov/sequence.html?DocID=20030099974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel isolated polypeptide associated with breast cancer, useful fodetecting presence of polypeptide in sample, as a marker for breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lillie J,
                                                                                                                                                                                                                                                                                                                                                       Human; cytosine methylation; 5'-CpG-3'; uracil; cytosine; diagnosis; drug; side effect; cancer; central nervous system; cardiovascular; gastrointestinal; respiratory system; single nucleotide polymorphism SNP; cell differentiation; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide
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                                                                                                             Olek A,
                                                                                                                                                                            01-SEP-2000;
05-SEP-2000;
                                                                                                                                                                                                                           01-SEP-2001; 2001WO-EP010074
                                                                                                                                                                                                                                                             07-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABQ41262;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABQ41262 standard;
                                                                                 WPI; 2002-371829/40.
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                                                                                                                                               (EPIG-)
                                                                                                                                                                                                                                                                                                                          sapiens.
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                                                                                                                                               EPIGENOMICS AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATGATGGCGGCGGAGTGTGA 114
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                                                                                                               Piepenbrock
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                              2000DE-01043826
2000DE-01044543
                                                                                                                                                                                                                                                                                                                                                                                                                                        for detecting cytosine methylation SEQ ID NO 27853.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA; 1276
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                                                                                                                 Berlin
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Pred. No.
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Determining the degree of cytosine diagnosis and prognosis, comprises from chemically treated DNA.

methylation in genomic DNA, selective hybridization of

useful for

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RESULT 9
ABQ41263/
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Best Local (
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ABQ41263
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                                                       Determining the degree of cytosine diagnosis and prognosis, comprises from chemically treated DNA.
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05-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                  drug; side effect; cancer; central nervous system; cardiovascular; gastrointestinal; respiratory system; single nucleotide polymorphi SNP; cell differentiation; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                               Human; cytosine methylation; 5'-CpG-3'; uracil; cytosine; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide for detecting cytosine methylation SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1276 BP;
                                                                                                                    WPI; 2002-371829/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATGATGGCGGCGGAGTGTGA
                                                                                                                                                 Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATGATGGCGGCGGAGTGTGA 601
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                                                                                                                                                                                                            2000DE-01043826.
2000DE-01044543.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      205 A; 151 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
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Pred. No. 0.18;
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                                                                       methylation in genomic DNA, useful for selective hybridization of amplicons
                                                                                                                                                  Guetig
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This invention describes a novel method for determining

the degree

of.

The invention relates to a novel colon specific protein (CSP) (I) and the nucleic acid (CSNA) encoding it. A CSP of the invention has cytostatic activity. The protein is useful as a vaccine, and the nucleic acid may have a use in gene therapy. A CSP is useful for determining the present

(CSP) (I) and the

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SEQ ID

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88;

547pp; English.

Claim 12; 56pp + Sequence Listing; 56pp; German

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RESULT 10
ADP46512/c
ID ADP465
XX ADP466
XX ADP4080
XX ADP465
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Best Local S
Matches 20
                                                                                                                                                                                                                                  Novel colon specific protein derived from normal and neoplastic colon cell, useful as vaccine in treating colon cancer and in identifying, diagnosing, monitoring, staging, imaging colon cancer and non-cancerou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-2002;
04-DEC-2002;
                                                                                                                                                                                                                disease state
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADP46512 standard; DNA; 1716
                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-480623/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DIAD-) DIADEXUS INC.
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2002US-0431144P.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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RESULT 11
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a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate call carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a patient; (ii) is also useful as a pharmacodyanamic or pharmaconic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of a colon specific protein in a sample. A CSNA is useful for determining the presence of a colon specific nucleic acid (CSNA) in a sample. The CSP and CSNA are useful for diagnosing or monitoring the presence and metastases of colon cancer in a patient. A method of the invention is useful for treating a patient with colon cancer. A CSP and CSNA are useful in identifying, diagnosing, monitoring, staging, imaging colon cancer and non-cancerous disease state in colon, and as a vaccine for treating colon cancer and non-cancerous disease states in colon. The present sequence represents a CSNA of the invention.
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25-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer.
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; 2000US-0189862P.
; 2000US-0207454P.
; 2000US-0211314P.
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marker; gene; ss.
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Pred. No. 0.18;
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                                                                                                                                                                                                                           The invention relates to a novel colon specific protein (CSP) (I) and the nucleic acid (CSNA) encoding it. A CSP of the invention has cytostatic activity. The protein is useful as a vaccine, and the nucleic acid may have a use in gene therapy. A CSP is useful for determining the presence of a colon specific protein in a sample. A CSNA is useful for determining the presence of a colon specific nucleic acid (CSNA) in a sample. The CSP and CSNA are useful for diagnosing or monitoring the presence and metastases of colon cancer in a patient. A method of the invention is useful for treating a patient with colon cancer. A CSP and CSNA are useful in identifying, diagnosing, monitoring, staging, imaging colon cancer and non-cancers state in colon, and as a vaccine for treating colon cancer and non-cancerous disease states in colon. The present sequence represents a CSNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ds; gene; human; colon specific protein; CSP; CSNA; cytostatic; vaccine;
gene therapy; colon specific nucleic acid; cancer; colon;
chromosome 2q37.1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel colon specific protein derived from normal and neoplastic colocell, useful as vaccine in treating colon cancer and in identifying, diagnosing, monitoring, staging, imaging colon cancer and non-cancer
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04-DEC-2002;
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2002US-0431144P.
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                                                                                                                                                                                        776 A; 418 C;
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                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         non-cancerous
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                                                                                                   Gaps
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ADR07433/c
                                                                                                                                                      CC CDNA clones obtained by an oligo-capping method, where none of these CC clones are identical to any known human mRNAs. The present invention CC describes an immunoassay to identify agonists and antagonists, as well as CC antibodies, antisense molecules and siRNAs that can all be used to bind to and modulate expression of the cDNA molecules. As such, these CC molecules are useful for diagnostic markers or therapeutic targets for the various diseases or morbid states. In particular, they are useful in CC gene therapy for treating osteoporosis, neurological disease, Alzheimer's CC disease, Parkinson's disease, dementia, short memory and various cancers, CC as well as for maintaining equilibrium of sense or motor function, and CC they exhibit osteopathic, neuroprotective, nootropic, antiparkinsonian, CC cytostatic and tranquiliser activities. This polynucleotide is a full CC length human cDNA sequence of the invention. NOTE: This sequence is not compared to the specification but can be obtained on CC CD-ROM from the European Patent Office, Vienna Sub-office.
                                                               Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isogai T,
Wakamatsu
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to novel, isolated full length human of molecules and the encoded proteins thereof. Specifically, it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New 1995 cDNA, useful Alzheimer's diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-FEB-2003; 2003JP-00102207.
09-MAY-2003; 2003JP-00131452.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP1447413-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Full length human cDNA useful for treating neurological disease Seq 939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADR07433;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parkinson's disease; dementia; short memory; cancer; sense or motor function; emotional reaction; fear response; panic; osteopathic; neuroprotective; nootropic; antiparkinsonian; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADR07433 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tranquiliser.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (REAS-) RES ASSOC BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-FEB-2004; 2004EP-00003145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2004-583265/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens.
 150
                                                                              Similarity
ATGATGGCGGCGGAGTGTGA 131
                                                                                                                              2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 939; 2686pp; English.
                              ATGATGGCGGCGGAGTGTGA
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                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                              B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             an; oligo-capping method; diagnostic
  neurological disease; Alzheimer's di
                                                                                                                             748 A; 413 C; 665 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ů,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                             100.0%; Score 20; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for treating osteoporosis, neurological diseases, Parkinson's diseases, dementia and various cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nishikawa T, Isono Y, Sugiyama Nagai K, Irie R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease; Alzheimer's disease;
                                                               0
                              20
                                                                 Mismatches
                                                                                                                             574 T; 0 U; 0 Other;
                                                                                            DB 13; Length 2400;
                                                                              .18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dementia and various cancers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                               refers to
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TV AX PX BX AX P F F X P X BX AX BX 
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                                                                                                                                                                                                                                              RESULT 14
                                                                                                                                                                                                                                                                                                                        This cDNA sequence codes for the P95 (or nucleolin) protein of the newly clidentified V3 loop HIV receptor. This novel protein complex receptor for CC HIV retroviruses consists of an association of 3 proteins named CC P95/nucleolin, P40/PHAPII and P30/PHAPI (see AAW84052-54). These proteins were isolated from human CD4+ CEM T-cell extracts using an affinity containing either the pseudopeptide S(KpsiCH2)NPR)-template assembled synthetic peptide or a synthetic V3 loop peptide (see CC AAW84055). P95 was identified as nucleolin by amino acid sequence analysis. The invention also concerns peptidic or non-peptidic molecules having the ability to alter and/or prevent the binding of the novel HIV receptor to the HIV retrovirus, and to pharmaceutical and diagnostic compositions containing such molecules. Methods are provided for screening for new active molecules, and to methods of screening special containing such molecules. Wethods are provided for screening for new active molecules and to methods of screening special that survive long-term HIV infection or who are HIV-resistant. Such
                                                                                                                   Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                          Sequence 2518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated V3 loop HIV receptor - comprises P95/nucleolin, P40/PHAPII and P30/PHAPI proteins, used to develop products for the treatment and prevention of HIV infection.
                                                                                                                                                                                                                                                                                                    genetically defective polynucleotides can be used in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 17; Fig 49(14); 267pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV71745 standard; cDNA; 2518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-PSDB; AAW84052.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9840480-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hovanessian A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INSP ) INST PASTEUR.
(CNRS ) CENT NAT RECH SCI.
113
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                                   ATGATGGCGGCGGAGTGTGA 20
   ATGATGGCGGCGGAGTGTGA 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Guichard
                                                                                                                   100.0%; ilarity 100.0%; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleolin; infection; therapy; diagnosis; ss
                                                                                                                                                                                                                                              BP; 823 A; 459 C; 711 G; 525 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                V3 loop; human immunodeficiency virus; retrovirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0040969P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98WO-EP001409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q
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                                                                                                                       Score 20; DB
Pred. No. 0.1
0; Mismatches
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                                                                                                                                                                                    DB
                                                                                                                                                       .18;
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                                                                                                                          Gaps
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RESULT 15 AAC92553/c ID AAC92553 standard; DNA; 2518

AAC92553;

27-MAR-2001

(first entry)

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cc known as P92 or C23) is the most abundant nucleolar phosphosprotein in CC actively growing cells. Nucleolin primarily participates in ribosome CC biogenesis and transport of ribosomes, to represent the components, being able to CC ribonucleopyrotein consensus sequence. However, it has also been shown to be involved in cytokinesis, nucleogenesis, cell proliferation and growth, cc chromatin decondensation. Nucleolin is a member of the Ag-NoR (active ribosomal gene located in the nucleolar organiser region) family of CC proteins which are markers of active ribosomal genes and whose CC presence of antibodies against nucleolin are associated with the prediction of tumour growth rate. The connective tissue diseases such as systemic lupus erythematosus (SLE) and cc scleroderma-like chronic graft versus host disease. The invention relates to antisense oligonucleotides targetted to the nucleolin gene, which can be connected to target different regions of the human nucleolin mRNA, can and were analysed for their effect on nucleolin mRNA levels by and seful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for diagnosis, prevention and treatment of conditions associated useful for the prevention and treatment of conditions associated useful for the prevention and treatment of conditions associated useful for the prevention and treatment of conditions associated useful for the prevention and treatment of conditions associated useful for the prevention and treatment of conditions associated useful for the prevention and the prevention and the prevention and the prevention and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel antisense compound targeted to human nucleolin which specifically hybridizes with and inhibits the expression of human nucleolin, useful for modulating the expression of nucleolin in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human nucleolin DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Col 47-52; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-079848/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents DNA encoding human nucleolin. Nucleolin (also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAB48964.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                              with nucleolin expression, such as tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-00433699
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /product= "Human nucleolin"
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                                                                                   formation,
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Query Match Best Local Similarity

100.0%;

Score 20; DB 4 Pred. No. 0.18;

DB 4;

Length 2518;

Sequence 2518 BP; 823 A; 459 C; 711 G; 525 T; 0 U; 0 Other;

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                                Matches
113 ATGATGGCGGCGGAGTGTGA
            1 ATGATGGCGGCGGAGTGTGA 20
                                20;
                                Conservative
                                 0
   94
                                 Mismatches
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Indels

0,

Gaps

0

Search completed: April 11, 2005, 23:56:49 Job time : 432 secs